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RAW SEQUENCE LISTING PATENT APPLICATION US/08/552,839

DATE: 02/12/96 TIME: 10:59:38

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INPUT SET: S8653.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING ENTER
2 3	(1) G	eneral Information:
4 5 6 7 8	(i)	APPLICANT: WANG, QING FINER, MITCHELL H. JIA, XIAO-CHI
9 10	(ii)	TITLE OF INVENTION: NOVEL ADENOVIRAL VECTORS, PACKAGING CELL LINES, RECOMBINANT ADENOVIRUSES, AND METHODS
11 12 13	(iii)	NUMBER OF SEQUENCES: 13
14 15 16 17 18 19 20 21	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: CELL GENESYS, INC. (B) STREET: 322 LAKESIDE DRIVE (C) CITY: FOSTER CITY (D) STATE: CALIFORNIA (E) COUNTRY: USA (F) ZIP: 94404
22 23 24 25 26 27	(V)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
28 29 30 31 32	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: (B) FILING DATE: 03-NOV-1995 (C) CLASSIFICATION:
33 34 35 36 37	(viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: KRUPEN, KAREN I. (B) REGISTRATION NUMBER: 34,647 (C) REFERENCE/DOCKET NUMBER: CELL 16.3
38 39 40 41 42	(ix)	TELECOMMUNICATION INFORMATION: (A) TELEPHONE: (415)358-9600 X131 (B) TELEFAX: (415)349-3792
43 44	(2) INFO	RMATION FOR SEQ ID NO:1:
45	(i)	SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31 base pairs

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47 48 49	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
50 51 52 53	(ii) MOLECULE TYPE: DNA (genomic)	
54 55 56	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
57 58	GCGCAAGCTT CGGGAGTGGG AGATAAGGCT C	31
59	(2) INFORMATION FOR SEQ ID NO:2:	
60	// GROVENGE GUADAGMEDICMIGG.	
61	(i) SEQUENCE CHARACTERISTICS:	
62	(A) LENGTH: 31 base pairs(B) TYPE: nucleic acid	
63		
64	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
65	(D) TOPOLOGY: linear	
66	(ii) MOLEGULE MUDE: DNA (gonomic)	
67	(ii) MOLECULE TYPE: DNA (genomic)	
68 69		
70		
71	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
72	(AI) DEGOTION DEDOCATION DE 15 NOVEL	
73	GGCCTCTAGA AGTTCACTTG CCCTGATGAC A	31
74	Occiding Norionalis Contament	
75	(2) INFORMATION FOR SEQ ID NO:3:	
76	(-)	
77	(i) SEQUENCE CHARACTERISTICS:	
78	(A) LENGTH: 18 base pairs	
79	(B) TYPE: nucleic acid	
80	(C) STRANDEDNESS: single	
81	(D) TOPOLOGY: linear	
82		
83	(ii) MOLECULE TYPE: DNA (genomic)	
84		
85		
86		
87	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
88		
89	GAGGACTAAG GATTGATT	18
90		
91	(2) INFORMATION FOR SEQ ID NO:4:	
92		
93	(i) SEQUENCE CHARACTERISTICS:	
94	(A) LENGTH: 18 base pairs	
95	(B) TYPE: nucleic acid	
96	(C) STRANDEDNESS: single	
97	(D) TOPOLOGY: linear	
98		
99	(ii) MOLECULE TYPE: DNA (genomic)	

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```
100
101
102
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
103
104
     CGTGAGATTT TGGATAAG
                                                                                 18
105
106
      (2) INFORMATION FOR SEQ ID NO:5:
107
108
109
           (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 12 base pairs
110
                (B) TYPE: nucleic acid
111
                (C) STRANDEDNESS: single
112
113
                (D) TOPOLOGY: linear
114
115
          (ii) MOLECULE TYPE: DNA (genomic)
116
117
118
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
119
120
     CCTACGCACG AC
                                                                                 12
121
122
123
     (2) INFORMATION FOR SEQ ID NO:6:
124
           (i) SEQUENCE CHARACTERISTICS:
125
                (A) LENGTH: 12 base pairs
126
127
                (B) TYPE: nucleic acid
128
                (C) STRANDEDNESS: single
129
                (D) TOPOLOGY: linear
130
131
          (ii) MOLECULE TYPE: DNA (genomic)
132
133
134
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
135
136
     TGTTTGGGTT AT
                                                                                 12
137
138
     (2) INFORMATION FOR SEQ ID NO:7:
139
140
           (i) SEQUENCE CHARACTERISTICS:
141
142
                (A) LENGTH: 15 base pairs
143
                (B) TYPE: nucleic acid
144
                (C) STRANDEDNESS: single
145
                (D) TOPOLOGY: linear
146
147
          (ii) MOLECULE TYPE: DNA (genomic)
148
149
150
151
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
152
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RAW SEQUENCE LISTING PATENT APPLICATION US/08/552,839

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153 154	GAC	CCGTGAGG ATACT			
155 156	(2)	INFORMATION FOR SEQ ID NO:8:			
157		(i) SEQUENCE CHARACTERISTICS:			
158		(A) LENGTH: 29 base pairs			
159		(B) TYPE: nucleic acid			
160		(C) STRANDEDNESS: single			
161		(D) TOPOLOGY: linear			
162		(5) 100 200 200 200			
163		(ii) MOLECULE TYPE: DNA (genomic)			
164		(-2,			
165					
166					
167		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:			
168		(AI) DEGOLACE DESCRIPTION DEG ES NOVO			
169	GCCI	AATCTAG AGCTTCAGGA AATATGACT	29		
170	0007	REIOTRO ROCITOROSE INTERIOR			
171	(2)	INFORMATION FOR SEQ ID NO:9:			
172	(-,	Internation for bug in the second			
173		(i) SEQUENCE CHARACTERISTICS:			
174		(A) LENGTH: 29 base pairs			
175		(B) TYPE: nucleic acid			
176		(C) STRANDEDNESS: single			
177		(D) TOPOLOGY: linear			
178		(b) forobodi. Hitedi			
179		(ii) MOLECULE TYPE: DNA (genomic)			
180		(II) MODECOLD III . DNA (GONOMIC)			
181					
182					
183		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:			
184		(AI) Digolikol Dibokii ilokk Dig ID kotot			
185	САТО	CTCTCGA GGGAGAAGTC CACGCCTAC	29		
186					
187	(2)	INFORMATION FOR SEQ ID NO:10:			
188	(-,	International Control of the Control			
189		(i) SEOUENCE CHARACTERISTICS:			
190		(A) LENGTH: 25 base pairs			
191		(B) TYPE: nucleic acid			
192		(C) STRANDEDNESS: single			
193		(D) TOPOLOGY: linear			
194		(5)			
195		(ii) MOLECULE TYPE: DNA (genomic)			
196		(,, (5,			
197					
198					
199		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:			
200		-			
201	TCCATTTCTA GATCGCTGC GGTTG 25				
202					
203	(2) INFORMATION FOR SEQ ID NO:11:				
204	-				
205		(i) SEQUENCE CHARACTERISTICS:			

RAW SEQUENCE LISTING PATENT APPLICATION US/08/552,839

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		INPUT SET: S8653.raw
206	(A) LENGTH: 18 base pairs	
207	(B) TYPE: nucleic acid	
208	(C) STRANDEDNESS: single	
209	(D) TOPOLOGY: linear	
210	(•, •••••••	
211	(ii) MOLECULE TYPE: DNA (genomic)	
212	(11) 110220022 11121 2111 (3011011120)	
213		
214		
215	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
216	(XI) DEGORNOE DESCRIPTION. DEG ID NO.II.	
217	ACGTGGTACT TGTCCATC	18
217	ACGIGGIACI IGICCAIC	10
219	(2) INFORMATION FOR SEQ ID NO:12:	
220	(2) INFORMATION FOR BEQ ID NO.12:	
221	(i) SEQUENCE CHARACTERISTICS:	
221	(A) LENGTH: 30 base pairs	
222	(B) TYPE: nucleic acid	
223	(C) STRANDEDNESS: single	
225	(D) TOPOLOGY: linear	
226	(ii) Not Dour E MUDE. DWA (comerie)	
227 228	(ii) MOLECULE TYPE: DNA (genomic)	
228 229		
229		
230	() CHOURNER DECERTION. CEO ID NO.12.	
231	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
232	TACTAACACT ACCCGCTGCT CTTGCTCTTG	30
233	TACTARCACT ACCOUNTS CITACTOTIC	30
235	(2) INFORMATION FOR SEQ ID NO:13:	
236	(2) INFORMATION FOR BEQ ID NO.13.	
237	(i) SEQUENCE CHARACTERISTICS:	
238	(A) LENGTH: 30 base pairs	
239	(B) TYPE: nucleic acid	
240	(C) STRANDEDNESS: single	
240	(D) TOPOLOGY: linear	
241	(D) TOPOLOGI: IIMeal	
242	/ii) MOLEGULE MUDE, DNA /gonomig)	
243	(ii) MOLECULE TYPE: DNA (genomic)	
244 245		
245 246		
	(wi) GEOLIENGE DECORPORAN, GEO. ID NO.12.	
247 248	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
248 249	TACTAACCTA GGACGCGGTC CCAGATGTTG	30
	INCIANCUTA GUNCGUGUTU CUNGATUTTU	30
250		

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